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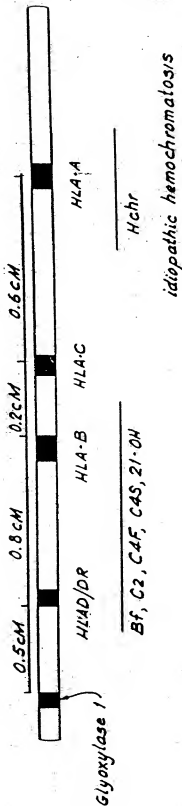


FIG. 1

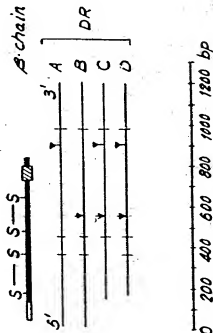


FIG. 4

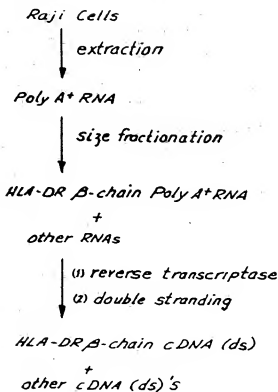
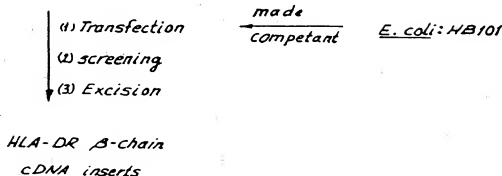
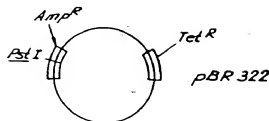
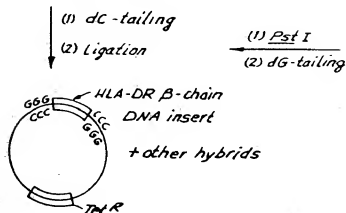


FIG. 2



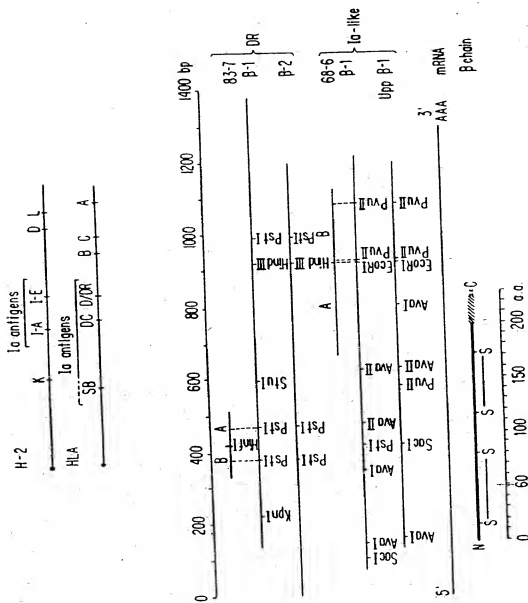


FIG. 3

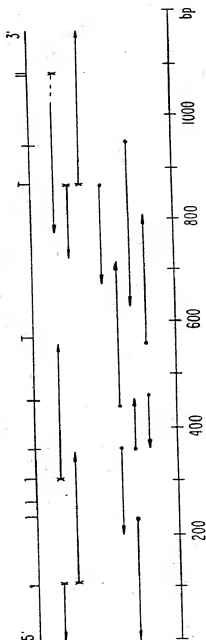


Applicant
Application No
Title
Attorney

Bernard F. Mach et al.
08/484,785
Confirmation No. 4894
DNA SEQUENCES CODING FOR THE DR β -CHAIN LOCUS OF
THE HUMAN LYMPHOCYTE ANTIGEN COMPLEX AND POLYPEPTIDES,
DIAGNOSTIC TYPING PROCESSES AND PRODUCTS RELATED THERETO
Margaret A. Piem, Reg. No. 30,709, Tel: (212) 596-9000

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Group Art Unit: 1634

FIG. 5A



DNA SEQUENCES CODING FOR THE DR B-CHAIN LOCUS OF
THE HUMAN LYMPHOCYTE ANTIGEN COMPLEX AND POLYPEPTIDES,
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-29
M V C L
ATG GTG TGT CTG 47

(G)_n CTCCTCTGGCCCTGGCTGCTCTCTCTCCAGC
K L P G G S S L A A L T V T 89
AAG CTC CCT GGA GGC TCC AGC TTG GCA GCG TTG ACA GTG ACA
L M V L S S R L A F A G D T 131
CTG ATG GTG CTG AGC TCC CGA CTG GCT TTC GCT GCG GAC ACC
R P R F L E L L K S E C H F 173
CGA CCA CGT TTC TTG GAG CTG CTT AAG TCT GAG TGT CAT TTC
F N G T E R V R F L E R H F 215
TTC AAT GGG ACG GAG CGG GTG CCG TTC CTG GAG AGA CAC TTC
H N Q E E Y A R F D S D V G 257
CAT AAC CAG GAG GAG TAC GCG CGC TTC GAC AGC GAC GTG GGG

FIG. 5B



50
E Y R A V R E L G R P D A E 299
GAG TAC CGG GCG GTG AGG GAG CTG GGG CGG CCT GAT GCC GAG

60
Y W N S Q K D L L E Q K R G 341
TAC TGG AAC AGC CAG AAG GAC CTC CTG GAG CAG AAG CGG GGC

80
Q V D N Y C R H N Y G V V E 383
CAG GTG GAC AAT TAC TGC AGA CAC AAC TAC GGG GTT GTG GAG

90
S F T V Q R R V H P Q V T V 425
AGC TTC ACA GTG CAG CGG CGA GTC CAT CCT CAG GTG ACT GTG

100
Y P A K T Q P L Q H H N L L 467
TAT CCT GCA AAG ACC CAG CCC CTG CAG CAC AAC CTC CTC

110
V C S V S G F Y P G S I E V 509
GTC TGC TCT GTG AGT GGT TTC TAT CCA GGC AGC ATT GAA GTC

FIG. 5C

DNA SEQUENCES CODING FOR THE DR B-CHAIN LOCUS OF
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220 G A G L F I Y F R N Q K G H
GGG GCC GGG CTG TTC ATC TAC TTC AGG AAT CAG AAA GGA CAC

230 S G L Q P T G F L S
TCT GGA CTT CAG CCA ACA GGA TTC CTG AGC TGA AGTGACAGATGA

237 CAATTAAAGGAAGATCTTCTTCCCCAGCTTTGCAGGATGAAAAGCTTTCCCGCC
TGGCTGTTATCTTCCACGAGAGAGGGCTTCTCAGGACCTAGTTGCTACTGGTT

803 CAGCAACTGCAGAAAATGTCTCCCTTGTGGCTTCTCCTCAGTTCCTGCCCTTGGCC
1012 TGAAGTCCCACCATTTGATGGCAGGCCTCATCTTCAAACCTTTGTGCTCCCCCTTG
1067 CCTAAACCCATATGGCCTCCTGTGCATCTGTACTACCTACCCCTGTACCAACAAACATTT
1122 ACATTATTAAATGTTTCTCAAAGATGGAGTTAAAAA(C)_n
1160

FIG. 5E



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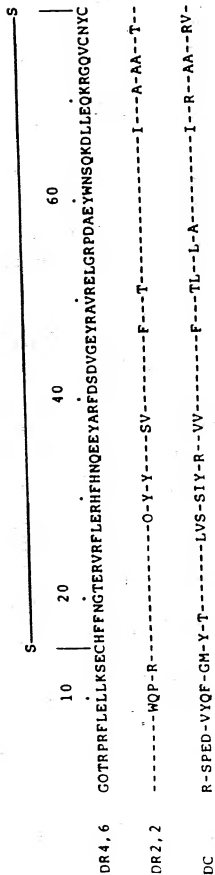


FIG. 6A



	80	
	100	
DR 4, 6	RHNYGVVESFTQRRVHPQVTVPFAKTPQLQHNLV	
DR 2, 2	-----Q-K-----S-----	
DC	-----QLELRT-L-----E-T--IS-SR-EA-N-----	

FIG. 6B

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Bernard F. Mach et al.
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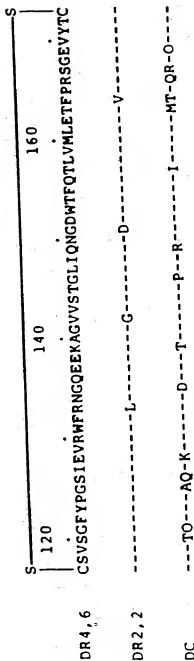


FIG. 6C

08/484,786
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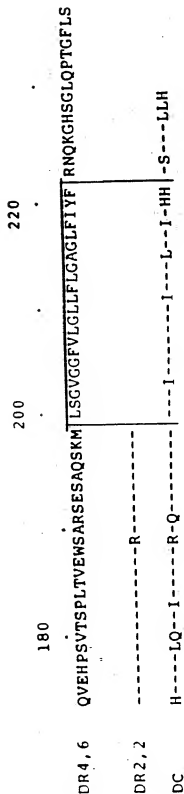


FIG. 6D



FIG. 7

[illegible]

FIG. 7A

¹⁶⁰
 I ACC TTC CAG ACC CTG GTG ATG CTG GAA ACA GAA GGT TAC ACC TGC CAA GTG GAG CAC CCA AGC CTG ACC AGC 694
¹⁷⁰
 P CCT CTC ACA GTG GAA TGG AGA GCA CGG TCT GAA TCT GCA CAG AGC AAG ATG CTG AGT GGA GTG GGG GGC TTC GTG CTG GGC CTG CTC 781
¹⁸⁰
 F TGC GAA GGC GGC CTG TTC ATC TAC TTT AGG AAT CAG AAA GGA CAC TCT GGA CTT CAG CCA ACA GGA TTC CTG AGC TGA AGT GAA 868
¹⁹⁰
 GAT GAC CAC ATT CAA GGA AGA ACC TTC TGC CCC AGC TTT GCA GGA TGA AAC ACT TCC CCG CTT GGC TCT CAT TCT TCC ACA AGA GAG 955
²⁰⁰
 ACC TTT CTC CGG ACC TGG TTG CTA CTG GTT CAG CAG CTC TGC AGA AAA TGT CCT CCC TTG TGG CTG CCT CAG CTC GTA CCT TTG GCC 1042
²¹⁰
 TGA AGT CCC AGC ATT AAT GGC AGC CCC TCA TCT TCC AAG TTT TGT GCT CCC CTT TAC CTA ATG CTT CCT GCG TCC CAT GCA TCT GTA 1129
²²⁰
 CTC CTG CTG TGC CAC AAA CAC ATT ACA TTA AAT GTT TCT CAA ACA TGG AGT TAA AAA AAA AAA AAA AAA AAA AA 1215
²³⁰
 GAT GAC CAC ATT CAA GGA AGA ACC TTC TGC CCC AGC TTT GCA GGA TGA AAC ACT TCC CCG CTT GGC TCT CAT TCT TCC ACA AGA GAG 955
²⁴⁰
 ACC TTT CTC CGG ACC TGG TTG CTA CTG GTT CAG CAG CTC TGC AGA AAA TGT CCT CCC TTG TGG CTG CCT CAG CTC GTA CCT TTG GCC 1042
²⁵⁰
 TGA AGT CCC AGC ATT AAT GGC AGC CCC TCA TCT TCC AAG TTT TGT GCT CCC CTT TAC CTA ATG CTT CCT GCG TCC CAT GCA TCT GTA 1129
²⁶⁰
 CTC CTG CTG TGC CAC AAA CAC ATT ACA TTA AAT GTT TCT CAA ACA TGG AGT TAA AAA AAA AAA AAA AAA AAA AA 1215

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FIG. 8

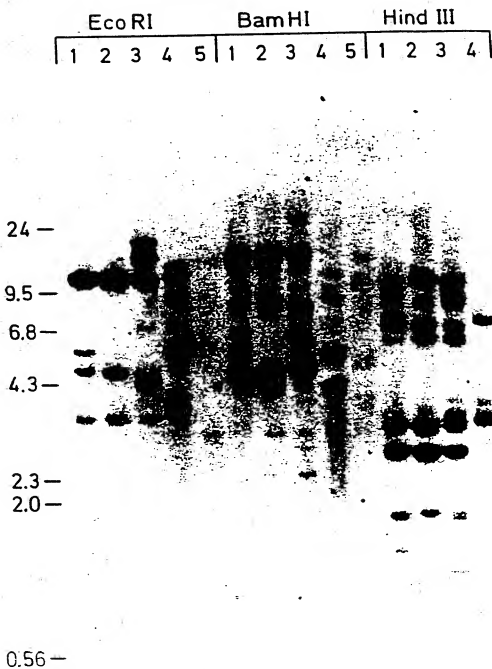




FIG. 9

Region I

	8	9	10	11	12	13	14
AA	L	E	L	L	K	S	E
HLA-DR- β -A	T	T	G	G	A	G	G
HLA-DR- β -	T	T	G	G	A	G	G
	L	E	Q	V	K	H	E

Region II

	26	27	28	29	30	31	32
AA	F	L	E	R	H	F	H
HLA-DR- β -A	T	T	C	G	A	C	A
HLA-DR- β -	T	T	C	G	A	C	A
	F	L	D	R	Y	F	Y

Region III

	72	73	74	75	76	77	78
AA	R	G	Q	V	D	N	Y
HLA-DR- β -A	C	G	G	C	A	A	T
HLA-DR- β -	C	G	G	C	A	A	T
	R	A	A	V	D	T	Y